

- G2
30. A method for designing primers, comprising the steps of:
- selecting at least one DNA nucleotide sequence from a genomic DNA database;
  - predicting a plurality of exons of said selected DNA nucleotide;
  - using each of the predicted exons as a template to design one corresponding primer pair for each of the predicted exons and for designing corresponding primer pairs for the predicted exons simultaneously; and
  - automatically collating said plurality of primer pairs with said predicted exons and the DNA nucleotide sequence.

- G3
34. A primer design system, comprising:
- means for selecting at least one genomic DNA nucleotide sequence from a database including a plurality of DNA nucleotide sequences;
  - means for predicting a plurality of exons of said selected DNA nucleotide and for storing positions of the predicted exons;
  - means for using each of the predicted exons as a template to design one corresponding primer pair for each of the predicted exons and for designing corresponding primer pairs for the predicted exons simultaneously; and
  - means for evaluating specificity of each designed primer or each designed primer pair.